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or bodily fluid from a normal human control, wherein an increase in the levels of the CSG determined in step (b) as compared to levels of the CSG measured in a sample of cells, tissues or bodily fluid from a normal human control is associated with a cancer that is progressing and a decrease in the determined levels of the CSG in the first sample as compared to the second sample is associated with a cancer that is regressing or in remission.

REMARKS

Claims 1 through 5 are pending in the instant application.

Claims 1 through 5 have been rejected. Claim 1-3 have been amended. No new matter has been added by these amendments.

Reconsideration is respectfully requested in light of these amendments and the following remarks.

I. Rejection of Claims 1-5 under 35 U.S.C. § 112, second paragraph

The rejection of claims 1-5 under 35 U.S.C. § 112, second paragraph, has been maintained, as the Examiner suggests that the recitation of CSG is indefinite. Accordingly, in an earnest to

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advance the prosecution of this case and in accordance with the Examiner's suggestion, Applicants have amended claim 1 to recite "colon specific gene (CSG)". Withdrawal of this rejection is therefore respectfully requested.

II. Rejection of Claims 1-5 under 35 U.S.C. § 112, first paragraph

Claims 1 through 5 have been rejected under 35 U.S.C. § 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor, at the time the application was filed, had possession of the claimed invention. Specifically, the Examiner suggests that Applicants did not point out in the amendment filed July 5, 2001, where support for amendments made to the claims can be found in the specification.

With respect to the phrases "a second sample of cells" and "first sample as compared to the second sample", it is respectfully pointed out that the claims 2 and 3 have been amended to delete these phrases. With respect to the phrase "comprising a polynucleotide sequence or its complement capable

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of hybridizing under stringent conditions with SEQ ID NO:1 or a polypeptide encoded thereby, support for this amendment is provided in the specification at page 3, lines 17-29, and page 7, lines 2 through 15.

Accordingly, withdrawal of this rejection is respectfully requested.

III. Rejection of Claims 1 and 2 under 35 U.S.C. § 102(b)

Claims 1 and 2 have been rejected under 35 U.S.C. 102(b) as being anticipated by U.S. Patent 5,733,748. Claims 1 and 2 have also been rejected under 35 U.S.C. § 102(b) as being anticipated by WO 96/39419. The Examiner suggests that these references disclose methods for diagnosing the presence of colon cancer and metastases of colon cancer which are based on determining the levels of a CSG comprising a polynucleotide sequence such as GCT (nucleic acid residues 9-11 and 13-15 of Figure 1 of these references) that would hybridize under stringent conditions with SEQ ID NO:1. Applicants respectfully traverse this rejection.

Contrary to the Examiner's suggestion, none of the 13 colon specific genes taught in the cited prior art references exhibit sufficient homology to the CSG of the present invention, namely

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SEQ ID NO:1, to hybridize under stringent conditions with SEQ ID NO:1. As evidence of this fact, Applicants are providing herewith a definition from the Life Sciences Dictionary of hybridization stringency. Hybridization stringency is defined as:

the percentage of nucleotides which must match on two unrelated single-stranded nucleic acid molecules before that will base pair with each other to form a duplex, given a certain set of physical and chemical conditions. . . . In general, if the percentage of matching nucleotides is lower than 70 percent, the two single-stranded nucleic acid molecules are considered nonhomologous and any hybridization is considered nonstringent.

Since the percentage of matching nucleotides of the colon specific genes taught in the cited prior art references and SEQ ID NO:1 of the instant invention is lower than 70 percent (see BLAST searches provided herewith), any hybridization occurring would be considered nonstringent by those of skill in the art. Accordingly, the prior art references do not teach a CSG comprising a polynucleotide sequence or its complement capable of hybridizing under stringent conditions with SEQ ID NO: 1 as claimed.

MPEP § 2131 is quite clear; to anticipate a claim, the reference must teach every element of the claim. Since the cited prior art references do not teach every element of the claims,

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they cannot anticipate the claimed invention. It is therefore respectfully requested that this rejection be withdrawn.

IV. Rejection of Claims 1-5 under 35 U.S.C. § 103(a)

Claims 1 through 5 have been rejected under 35 U.S.C.

103(a) as being unpatentable over U.S. Patent 5,733,748 and WO

96/39419. The Examiner suggests that it would have been prima

facie obvious to one of ordinary skill in the art at the time the

claimed invention was made to implement the methods of staging

and monitoring colon cancer in a patient for changes in staging

as well as for the onset of metastasis considering the methods of

diagnosing the presence of colon cancer and metastases have been

well established. Applicants respectfully traverse this

rejection.

MPEP 2143 is clear; to establish a prima facie case of obviousness invention, three basic criteria must be met. First, there must be some suggestion or motivation, either in the references themselves or in the knowledge generally available to one of ordinary skill in the art, to modify the reference or to combine the reference teachings. Second, there must be a reasonable expectation of success. Finally, the prior art

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reference must teach or suggest all the claim limitations.

As discussed in Section III, supra, neither of the cited prior art references teach the CSG of SEQ ID NO:1 nor a sequence that would hybridize under stringent conditions with SEQ ID NO:1. Nor is there any suggestion in these references of a CSG of SEQ ID NO:1 or a sequence that would hybridize under stringent conditions with SEQ ID NO:1. Accordingly, these references do not teach or suggest all the claim limitations. These references also provide no reasonable expectation of success that a CSG of SEQ ID NO:1 or a sequence that would hybridize under stringent conditions with SEQ ID NO:1 would be useful in diagnosing, monitoring staging and imaging cancer. There is also no motivation or suggestion provided in these references to modify their teachings to use a CSG of SEQ ID NO:1 or a sequence that would hybridize under stringent conditions with SEQ ID NO:1 to diagnose, monitor, stage or image cancer. Thus, the cited combination of references cannot render obvious the instant claimed invention.

Withdrawal of this rejection under 35 U.S.C. § 103(a) is therefore respectfully requested.

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V. Supplemental IDS

A Supplemental Information Disclsoure Statement is being provided herewith for consideration by the Examiner.

VI. Conclusion

Applicants believe that the foregoing comprises a full and complete response to the Office Action of record. Accordingly, favorable reconsideration and subsequent allowance of the pending claims is earnestly solicited.

Attached hereto is a marked-up version of the changes made to the specification and claims by the current amendment. The attached page is captioned "Version with Markings to Show Changes Made."

Respectfully submitted,

Sarinassylican

Jane Massey Licata Registration No. 32,257

Date: **January 25, 2001**

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VERSION WITH MARKINGS TO SHOW CHANGES MADE

In the Claims:

Please amend the claims as follows:

- 1. (amended) A method for diagnosing the presence of colon cancer in a patient comprising:
- (a) determining levels of a <u>colon specific gene(CSG)</u> comprising a polynucleotide sequence or its complement capable of hybridizing under stringent conditions with SEQ ID NO: 1, or a polypeptide encoded thereby, in cells, tissues or bodily fluids in a patient; and
- (b) comparing the determined levels of the CSG with levels of the CSG in cells, tissues or bodily fluids measured in a normal human control, wherein a change in determined levels of the CSG in said patient versus levels of the CSG measured in a normal human control is associated with the presence of colon cancer.
- 2. (amended) A method of diagnosing metastases of colon cancer in a patient comprising:
- (a) identifying a patient having colon cancer that is not known to have metastasized;
 - (b) determining levels of a CSG comprising a

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polynucleotide sequence or its complement capable of hybridizing under stringent conditions with SEQ ID NO: 1, or a polypeptide encoded thereby, in a first sample of cells, tissues or bodily fluid from said patient; and

- (c) comparing the determined levels of the CSG determined in step (b) with levels of the CSG measured in a second sample of cells, tissues or bodily fluid from a normal human control, wherein an increase in determined levels of the CSG in the first sample determined in step (b) as compared to the second sample levels of the CSG measured in a sample of cells, tissues or bodily fluid from a normal human control is associated with a cancer that has metastasized.
- 3. (amended) A method of staging colon cancer in a patient having colon cancer comprising:
 - (a) identifying a patient having colon cancer;
- (b) determining levels of a CSG comprising a polynucleotide sequence or its complement capable of hybridizing under stringent conditions with SEQ ID NO: 1, or a polypeptide encoded thereby, in a first sample of cells, tissues or bodily fluid from said patient; and
- (c) comparing the determined levels of the CSG determined in step (b) with levels of the CSG measured in a

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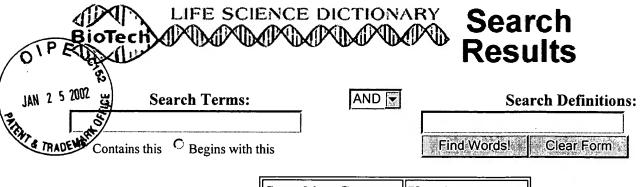
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second sample of cells, tissues or bodily fluid from a normal human control, wherein an increase in the determined levels of the CSG in the first sample determined in step (b) as compared to the second sample levels of the CSG measured in a sample of cells, tissues or bodily fluid from a normal human control is associated with a cancer that is progressing and a decrease in the determined levels of the CSG in the first sample as compared to the second sample is associated with a cancer that is regressing or in remission.



Searching Category	User input query
Searched Word	hybridization
Number of Results	14

1. 1. colony hybridization

Definition:

A genetics lab technique used to identify which colonies of bacteria on an agar plate contain a particular sequence of <u>DNA</u> or a particular gene. The technique involves pressing a nylon or nitrocellulose membrane onto the plate so that each colony contributes a small smudge of itself to the membrane, then treating the membrane with chemicals and heat, then washing the membrane with a labeled probe to find the specific DNA sequence. The smudges which are indicated by the <u>probe</u> are then compared back to the colonies on the agar plate. This technique is often used in conjunction with experiments involving the making of genomic libraries.

2. competition hybridization

Definition:

A lab technique used to determine how similar two strands of single-stranded <u>nucleic acids</u> are to each other by putting them with a third strand (called a standard) and observing how well they can bond with each other to become double-stranded (how well they <u>hybridize</u>).

3. cross-hybridization (cross hybridization)

Author: Susan A. Hagedorn

Definition:

The <u>hydrogen bonding</u> of a single-stranded <u>DNA</u> sequence that is partially but not entirely <u>complementary</u> to a single-stranded <u>substrate</u>. Often, this involves <u>hybridizing</u> a DNA <u>probe</u> for a specific DNA <u>sequence</u> to the <u>homologous</u> sequences of different <u>species</u>.

4. DNA hybridization

Definition:

A lab technique used to find out how closely related two or more separate strands of <u>DNA</u> from different <u>species</u> are to each other. The technique involves radioactive labeling.

5. DNA-RNA hybridization

Definition:

A type of <u>hybridization</u>. In this case, a strand of <u>DNA</u> is joined with a <u>complementary</u> strand of <u>RNA</u> to form a double-stranded molecule (or one which is partly double-stranded, if one of the original single strands is shorter than the other).

6. FISH (fluorescence in situ hybridization)

Definition:

A <u>physical mapping</u> approach that uses <u>fluorescent</u> tags to detect <u>hybridization</u> of <u>probes</u> with <u>metaphase chromosomes</u> and with the less-condensed <u>somatic interphase</u> chromatin.

7. hybridization

Definition:

- 1. The process of joining two <u>complementary</u> strands of <u>DNA</u> or one each of DNA and RNA to form a double- stranded molecule.
- 2. The mating of individuals from different species or sub-species.

8. hybridization stringency

Definition:

The percentage of <u>nucleotides</u> which must match on two unrelated single-stranded <u>nucleic acid</u> molecules before they will <u>base pair</u> with each other to form a <u>duplex</u>, given a certain set of physical and chemical conditions. The hybridization stringency is used to determine when a <u>hybridization probe</u> and a target nucleic acid will come together, and can be set by the researcher by varying the conditions. In general, if the percentage of matching nucleotides is lower than 70 percent, the two single-stranded nucleic acid molecules are considered <u>nonhomologous</u> and any <u>hybridization</u> is considered <u>nonstringent</u>.

9. in situ hybridization

Definition:

Use of a <u>DNA</u> or <u>RNA</u> probe to detect the presence of the <u>complementary DNA</u> sequence in cloned bacterial or cultured eukaryotic cells.

Also used for locating genes on chromosomes. The process is:

- 1. Prepare microscope slide with cells in metaphase of mitosis.
- 2. Treat slide with a weak base. Thus denaturing the DNA.
- 3. Pour radioactively labeled probe onto the slide.
- 4. Expose slide to photographic emulsion for a few days or weeks.
- 5. Develop emulsion.

10. introgressive hybridization

Definition:

The incorporation into a population's gene pool of genes from a different species.

11. Northern blot (Northern hybridization, Northern blotting) Definition:

A technique similar to <u>Southern blotting</u>, though it is used for <u>RNA</u>. In this technique, RNA fragments are transferred from an <u>agarose</u> gel to a nitrocellulose filter, where the RNA is then <u>hybridized</u> to a radioactive <u>probe</u>.

12. probe (hybridization probe)

Definition:

A single-stranded <u>nucleic acid</u> molecule with a known <u>nucleotide</u> sequence which is <u>labeled</u> in some way (for example, <u>radioactively</u>, <u>fluorescently</u>, or immunologically) and used to find and mark certain <u>DNA</u> or <u>RNA</u> sequences of interest to a researcher by <u>hybridizing</u> to it.

13. Southern blot (Southern hybridization, Southern blotting) Definition:

A technique used for searching for a specific <u>DNA</u> fragment. The process is as follows:

- 1. Separate DNA fragments by gel electrophoresis
- 2. change pH of gel to basic, thus allowing disruption of H-bonds
- 3. blot gel with nitrocellulose paper
- 4. heat paper so as to fix DNA fragments
- 5. probe with labeled messenger RNA or cDNA
- 6. wash
- 7. complementary mRNA/cDNA fragments will have hybridized.

14. Western blot (Western hybridization, Western blotting) Definition:

A technique similar to Southern blotting, though it is used for proteins.

END

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BLASTN 2.2.1 [Apr

#DEX0075_1_vs_wo9639419 blastn

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PADENT
Reference: Altschul, Stephen F., Thomas L. Madden, Alejandro A. Schaffer,
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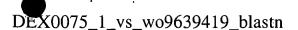
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wo9639419.nt



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GTACATACCTGGCTGGATTGACAATGATGAAATACAATGGAATCCACCAAGACCTGAAAT TAATAAGGATGATGTTCAACACAAGCAAGTGTGTTTCAGCAGAACATCCTCGGGAGGCTC ATTTGTGGCTTCTGATGTCCCAAATGCTCCCATACCTGATCTCTTCCCACCTGGCCAAAT CACCGACCTGAAGGCGGAAATTCACGGGGGCAGTCTCATTAATCTGACTTGGACAGCTCC TGGGGATGATTATGACCATGGAACAGCTCACAAGTATATCATTCGAATAAGTACAAGTAT TCTTGATCTCAGAGACAAGTTCAATGAATCTCTTCAAGTGAATACTACTGCTCTCATCCC AAAGGAAGCCAACTCTGAGGAAGTCTTTTTGTTTAAACCAGAAAACATTACTTTTGAAAA TGGCACAGATCTTTCATTGCTATTCAGGCTGTTGATAAGGTCGATCTGAAATCAGAAAT ATCCAACATTGCACGAGTATCTTTGTTTATTCCTCCACAGACTCCGCCAGAGACACCTAG TCCTGATGAAACGTCTGCTCCTTGTGCCTAATATTCATATCAACAGCACCATTCCTGGCA TTCACATTTTAAAAATTATGTGGAAGTGGGTAGGAGAACTGCAGTTGTCAATAGNCTAGG GGTGAATTTTTGTGCGGTGAATAAATAATSATTTCANCCTTTTTTTGRTTTATAAAAAAA CGGNTNCCCATTGGGNNTNTNGNGGGGGGGGNNTTTTAA >AAT45885 WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG6 cDNA partial clone. [Homo sapiens.] ${\tt AGTCGCTCTCTAGCCCTTCTCTGTGCCTCACCCTCTGGCAATGCCATTCAGGCCAGGTC}$ ${\tt TTCCTCCTATAGTGGAGAGTATGGAGGTGGTGGTAGAAGCGATTCTCTCATTCTGGCAA}$ CCAGTTGGACGCCCCATCACCGCCCTCCGGGTCCGAGTCAACACATACTACATCGTAGG GGAGGAGATCTTTCTGCACCCTGGGGAATCAGTGATCCAGGTTTCTGGGAAGTACAAGTG GTACCTGAAGAAGCTGGTATTTGTGACAGACAAGGGCCGCTATCTGTCTTTTGGGAAAGA CAGTGGCACAAGTTTCAATGCCGTCCCCTTGCACCCCAACACCGTGCTCCGCTTCATCAG TGGCCGGTCTGGTTCTCTCATCGATGCCATTGGCCTGCACTGGGATGTTTACCCCACTAG $\tt CTGCAGCAGATGCTGAGCCTCCTCTCTTTGGCAGGGGCACTGTGATGAGGAGTAAGAACT$ CCTTATCACTAACCCCCATC >AAT45887 WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG8 cDNA full-length clon e. [Homo sapiens.] CAGGAGGAGAGCCTTCCCCAAGCAAACAATCCAGAGCAGCTGTGCAAACAACGGTGCAT AAATAAGGCCTCCTGGACCATGAATGCGAGTCCGCTGAGCTGCGTACCGGAGCCCACGGT GGTCATGGCTGCCAGAGCGCTCTGCATGCTGGGGCTGGTCCTGGCCTTGCTGCTCCCAG GGTGGACTGCGGCTACCCCCATGTCACCCCCAAGGAGTGCAACAACCGGGGCTGCTGCTT TGACTCCAGGATCCCTGGAGTGCCTTGGTGTTTCAAGCCCCTGACAGGGAAGCAGGAATG CACCTTCTGAGGCACCTCCAGCTGCCCCCGGCGGGGGATGCGAGGCTCGGAGCACCCT TGCCCGGCTGTGATTGCTGCCAGGCACTGTTCATCTCAGCTTTTCTGTCCCTTTGCTCCC GGAAGCGCTTCTGCTGAAAGTTCATATCTGGAGCCTGATGTTTAACGTAGTCCCATGCTC САСССБААААААААААААААААААА >AAT45888 W09639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG9 cDNA full-length clon e. (Homo sapiens.) AAGCTCTTCTCACAGGACCAGCCACTAGCGCAGCTCGAGCGATGGCCTATGTCCCCGCAC TCAACGTGGGAATGTCTGTTTACATCCAAGGAGTGGCCAGCGAGCACATGAAGCGGTTCT ${\tt TCGTGAACTTTGTGGTTGGGCAGGATCCGGGCTCAGACGTCGCCTTCCACTTCAATCCGC}$ GGTTTGACGGCTGGGACAAGGTGGTCTTCAACACGTTGCAGGGCGGGAAGTGGGGCAGCG AGGAGGAGGAGGAGCATGCCCTTCAAAAAGGGTGCCGCCTTTGAGCTGGTCTTCATAGTCCTGGCTGAGCACTACAAGGTGGTGGTAAATGGAAATCCCTTCTATGAGTACGGGCACC TCAACTTCATCGGAGGCCAGCCCCTCCGGCCCCAGGGACCCCCGATGATGCCACCTTACC CTGGTCCCGGACATTGCCATCAACAGCTGAACAGCCTGCCCACCATGGAAGGACCCCCAA CCTTCAACCCGCCTGTGCCATATTTCGGGAGGCTGCAAGGAGGGCTCACAGCTCGAAGAA CCATCATCAAGGGCTATGTGCCTCCCACAGGCAAGAGCTTTGCTATCAACTTCAAGG TGGGCTCCTCAGGGGACATAGCTCTGCACATTAATCCCCGCATGGGCAACGGTACCGTGG TCCGGAACAGCCTTCTGAATGGCTCGTGGGGATCCGAGGAGAAGAAGATCACCCACAACC CATTTGGTCCCGGACAGTTCTTTGATCTGTCCATTCGCTGTGGCTTGGATCGCTTCAAGG TTTACGCCAATGGCCAGCACCTCTTTGACTTTGCCCATCGCCTCTCGGCCTTCCAGAGGG TGGACACATTGGAAATCCAGGGTGATGTCACCTTGTCCTATGTCCAGATCTAATCTATTC CTGGGGCCATAACTCATGGGAAAACAGAATTATCCCCTAGGACTCCTTTCTAAGCCCCTA >AAT45889 W09639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG10 cDNA partial clone. [Homo sapiens.] GTTGATATTAAAACCAGTGAAACCAAACATGACACCTCTCTGAAACCTATTAGTGTCTCC ${\tt GAGGACAACGATAACCGATCAGTGCTGAAAGGTGGTCCTTTCTCTGACAGCTACAGGCTC}$ TTTCAGTTCCATTTTCACTGGGGCAGTACAAATGAGCATGGTTCAGAACATACAGTGGAT GGAGTCAAATATTCTGCCGAGCTTCACGTGGCTCACTGGAATTCTGCAAAGTACTCCAGC CTTGCTGAAGCTGCCTCAAAGGCTGATGGTTTTGGCAGTTATTGGTGTTTTTGATGAAGGTT GGTGAGGCCAACCCAAAGCTGCAGAAAGTACTTGATGCCCTCCAAGCAATTAAAACCAAG TTCTGGACCTACCCTGGCTCTCTGACTCATCCTCTCTTTATGAGAGTGTAACTTGGATC ATCTGTAAGGAGAGCATCAGTGTCAGTTCAGAGCAGTTGGCACAATTCCGGAGCCTTCTA TCAAT WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG12 cDNA partial clone. >AAT45891 [Homo sapiens.] GTGGCAGAAGAAGATAGGTTGGAGACAATTGATTGCTCGATGATATAAAATGTTAAGTA CCATGAATGNATGCTGTTAGGCTGGAATGCGCCAAGATAAAAGGTGGGGCATGGCATCAA AAGGTAGGTCAACATATTAAATAATTCCATGTATTGAAATATCCAGAAAATATATAGACA GATCTATAGAGATAGAAACTGGTCTGCCCAGGACTAGGGGTTGTCTAAGGATAAGGAGCT TCTTTTTTGGATGGTGAAATAACCTAAAATATATTGTGCCATTGTTTGCACAACTTTGTG GAATATATTAAAAACCGGTTAATTGTACTCACTAAAATGTCCTCCTTCTTAAATTTAAGC TGTTTNCTGGACAAGAAAAGGGAAAGNNACCAAGGGGNAAAAATTTT

PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG13 cDNA partial clone.

[Homo sapiens.] GCCCTGGGCTTTGGGGGGGTCCCAAACATGGTATGCAGAAATGTGATGGTTACAGGTCAG TACAACCTCAGTCCTTAGAACCCCTCCACACTTCAGCTCTGCACCCACTTTCCTGTCATT

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>AAT45892



wo9639419.nt



 ${\tt CCATTCTCTTTAAAGGGGTTTTTATGGGGGCGCTTACTGCAGGTTAAAAATTGGGGNCCAC}$ >AAT45880 WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG1 partial cDNA clone. [Homo sapiens.} GCCAGGCAGCTGCCSACCAGGCCGTGTATGTGAAGGTCAAGGCTGAAGCCCGGGAA CTGCTGGGCCACCCGTGTCTCTGTGTCCTGTCTGTGGGTGCCAACTCACCACCTTTGAT ${\tt GGGGCCCGTGGTGCCACCACTCTCCTGGTGTCTATGAAGCTCTCTTCCCGCTGCCCAGGA}$ $\tt CTACAGAATACCATCCCCTGGTACCGTGTAGTTGCCGAAGTCCAGATCTGCCATGGCAAA$ ${\tt ACGGAGGCTGTGGGCCAGGTCCACATCTTCTTCCAGGATGGGATGGTGACGTTGACTCCA}$ AACAAGGGTGTGTGGGTGAATGGTCTCCGAGTGGATCTCCCAGCTGAGAAGTTAGCATCT GTGTCCGTGAGTCGTACACCTGATGGCTCCCTGCTAGTCCGCCAGAAGGCAGGGGTCCAG GTGTGGCTTGGAGCCAATGGGAAGGTGGCTGTGATTGTGAGCAATGACCATGCTGGGAAA AGGAGAAGCCAGCGATTGGGGAAWTGGAGAGCGCAGGGACTTTCTYCCMCATGTTAATGG GCTTGWTCCAGTTCATCCCACCAGGAACGAAGGATTTT WO9639419-A1 >AAT45881 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG2 fragment. [Homo sapi ens.l CAGGACTGCGTGTGCACGGACAAGGTGGACAACAACACCCTGCTCAACGTCATCGCCTGC ACCCACGTGCCCTGCAACACCTCCTGCAGCCCTGGGTTCGAACTCATGGAGGCCCCCGGG GAGTGCTGTAAGAAGTGTGAACAGACGCACTGTATCATCAAACGGCCCGACAACCAGCAC GTCATCCTGAAGCCCGGGGACTTCAAGAGCGACCCGAAGAACAACTGCACATTCTTCAGC TGCGTGAAGATCCACAACCAGCTCATCTCGTCCGTTTCCAACATCACCTGCCCCAACTTT ${\tt GATGCCAGCATTTGCATCCCGGGCTCCATCACATTCATGCCCAATGGATGCTGCAAGACC}$ TGCACCCTCGCAATGAGACCAGGGTGCCCTGCTCCACCGTCCCCGTCACCACGGAGGTT TCGTACGCCGGCTGCACCAAGACCGTCCTCATGAATCATTGCTCCGGGTCCTGCGGGACA ${\tt TTTGTCATGTACTCGGCCAAGGCCCAGGCCCTGGACCACAGCTGCTCCTGCAAAGAG}$ GAGAAAACCAGCCAGCGTGAGGTGGTCCTGAGCTGCCCCAATGGCGGCTCGCTGACACAC ACCTACACCCACATCGAGAGCTGCCAGTGCCAGGACACCGTCTGCGGGCTCCCCACCGGC ${\tt ACCTCCCGCCGGGCCCGGCGTTCCCCTAGGCATCTGGGGAGCGGGTGAGCGGGTGGGCA}$ ${\tt CAGCCCCTTCACTGCCCTCGACAGCTTTACCTCCCCCGGACCCTCTGAGCCTCCTAAGCT}$ ${\tt CGGCTTCCTCTTCAGATATTTATTGTCTGAGTTTTTGTTCAGTCCTTGCTTTCCAATA}$ ATAAACTCAGGGGGACATGCAAAAAAAAAAAAA >AAT45883 W09639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG4 cDNA partial clone. [Homo sapiens.] ATGAGTCCTGTGAAAAACAATGTGGGCAGAGGCCTAAACATCGCCCTGGTGAATGGAACC ACGGGAGCTGTGCTGGGACAGAAGGCATTTGACATGTACTCTGGAGATGTTATGCACCTA GTGAAATTCCTTAAAGAAATTCCGGGGGGTGCACTGGTGCTGGTGGCCTCCTACGACGAT ${\tt CCAGGGACCAAAATGAACGATGAAAGCAGGAAACTCTTCTCTGACTTGGGCAGTTCCTAC}$ GCAAAACAACTGGGCTTCCGGGACAGCTGGGTCTTCATAGGAGCCCAAAGACCTCAGGGGT CCAGAGCTGCTGGAGATGGAGGGCTGCATGCCCCCGAAGCCATTTTAGGGTGGCTGTGGC TCTTCCTCAGCCAGGGGCCTGAAGAAGYTCCTGCCTGCATTAGGAGTCANAGCCCGGCAG GCTGNAGGAGGAGCAGGGGGTGCTGCGTGGAAGGTGCTGCAGGCCTTGCACGCTGTG TCGCGCCT >AAT45886 WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG7 cDNA partial clone. [Homo sapiens.] ATTCTCAAATTTTATCTAAATGGATAAACTATTAACATAGAACATAAACCCCAATTCTCC ATTTCATTTTCTCTTAGGCATGAATCATACAAAACTCAATATAGAGCAATGTTTGTAAT GAATTGTTCTATTAACAAAGAGGAGGTTCTAAGATATAAAGCCTCAGAGAACAGGAAGAA AAGGCGGGTCCATAAGAAGATGAGGTCTAACCGGGAAGATGCTGCTGAGAAGGCAGAGAC AGATGTGGAAGAAATCTATCACCCAGTCATGTGCACTGAATGTTCCACTGAAGTGGCAGT TTACGACAAGGATGAAGTCTTTCATTTTTCAATGTTTTAGCAAGCCATTCCTAAACAGC CCAACTGGCATTTAATTACCCAATACTGTATATAAGGCAAATATGGACAGTTACTTTCCT CTTGCCTGTTCATATCCTTCAGTGACATTGAGGAAGCAGTGTTTCTCTTTTTAAAGGGGA ATTTTCCCCCTTATTGATGGGACTGATATTCATTCTGTTTTTGATGAACATTTGGAAACT GTCGGGCTTTTTATTAAAGCTCTGTAGAATTAAAATGTTCTGGAATTAT >AAT45890 WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG11 cDNA partial clone. [Homo sapiens.] CGGCTCCGGGCGGGCGTGGCCAGTGACTAGAAGGCGAGGCGCCGCGGGACCATGGCGGCG GCGGCGGACGAGCGAGTCCAGAGGCGAGAAGACGAGGAAGAGGAGCAGTTGGTTCT GGTTTTGGGCATTGACACTGAGAGGCCCATTCTGGCAATGGACAGCTGTGTCTTTGCTGG GGAGTATGAAGACACTCTAGGGACCTGTGTTATATTTGAAGAAAATGTTGAACATGCTGA TACAGAAGGCAATAATAAAACAGTGCTAAAATATAAATGCCATACAATGAAGAAGCTCAG CATGACAAGAACTCTCCTGACAGAGAAGAAGGAAGGAAGAAAACATAGGTGGGGTGGA ATGGCTGCAAATAAGGATATGGTTTCTCCCTTTGACCCAACAGGTTTGTTAACTTTTCTA CCATGAAATTGAGGACGAGGAAGTGGTAGCTTTCAGCCCCGTTAAATCTTTGGATTTGGG AGGGGGTGGGGTTTCAATG >AAT45882 WO9639419-A1 PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Human colon specific gene CSG3 fragment. [Homo sapi







DEX0075 1_vs_us5733748 blastn

E

0.29

0.29

4.6

4.6

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Reference: Altschul, Stephen F
                              ., Thomas L. Madden, Alejandro A. Schaffer,
Jinghui Zhang, Zheng Zhang, Webb Miller, and David J. Lipman (1997),
*Gapped BLAST and PSI-BLAST: a new generation of protein database search
programs*, Nucleic Acids Res. 25:3389-3402.
Query= DEX0075_1
         (2609 letters)
Database: us5733748.nt
          15 sequences; 9317 total letters
Searching.done
                                                                  Score
Sequences producing significant alignments:
                                                                   (bits) Value
AAV16668 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1...
AAV16676 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1...
                                                                       26
AAV16669 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1...
                                                                       24 1.2
                                      PR 06-JUN-1995 PF 06-JUN-1...
AAV16680 US5733748-A PA (HUMA-) HUMAN
                                                                       22 4.6
AAV16677 US5733748-A PA (HUMA-) HUMAN
                                      PR 06-JUN-1995 PF 06-JUN-1...
                                                                       22
AAV16674 US5733748-A PA (HUMA-) HUMAN
                                      PR 06-JUN-1995 PF 06-JUN-1...
                                                                       22
AAV16671 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1...
                                                                       22 4.6
           US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF
          06-JUN-1995 Polynucleotide sequence of a colon-specific
          gene. [Homo sapiens.]
         Length = 638
 Score = 26.3 bits (13), Expect = 0.29
 Identities = 13/13 (100%)
 Strand = Plus / Plus
Query: 825 actcaccaccttt 837
           1111111111111
Sbjct: 105 actcaccaccttt 117
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 606 gatggctccct 616
           ĬĦĬĦĦ
Sbjct: 382 gatggctccct 392
>AAV16676
```

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US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF
 06-JUN-1995 Polynucleotide sequence of a colon-specific
 gene. [Homo sapiens.]
Length = 1121
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Score = 26.3 bits (13), Expect = 0.29Identities = 13/13 (100%) Strand = Plus / Plus

Query: 1163 ggaagtggggcag 1175 Sbjct: 286 ggaagtggggcag 298

Score = 26.3 bits (13), Expect = 0.29Identities = 13/13 (100%) Strand = Plus / Minus

Query: 1339 ggattgaagtgga 1351 111111111111 Sbjct: 238 ggattgaagtgga 226

Score = 24.3 bits (12), Expect = 1.2Identities = 12/12 (100%) Strand = Plus / Minus

Query: 990 tttgaagggcat 1001 Sbjct: 329 tttgaagggcat 318



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06-JUN-1995 Polynucleotide sequence of a colon-specific
           gene. [Homo sapiens.]
          Length = 874
 Score = 24.3 bits (12), Expect = 1.2
 Identities = 12/12 (100%)
 Strand = Plus / Minus
Query: 69 gtaggtgtgtgt 80
           ĪĦĪĬĦĬĦĬĦ
Sbjct: 606 gtaggtgtgtgt 595
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Minus
Query: 1001 tcccctgagt 1011
            1111111
Sbjct: 855 tcccctgagt 845
Score = 22.3 bits (11), Expect = 4.6 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 2519 agtgtgaacag 2529
            Sbjct: 134 agtgtgaacag 144
>AAV16680
           US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF
           06-JUN-1995 Polynucleotide sequence of a colon-specific
           gene. (Homo sapiens.)
          Length = 600
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 620 ctccattctct 630
           11111111111
Sbjct: 479 ctccattctct 489
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 1140 atgaacaagcg 1150
            1111111111
Sbjct: 319 atgaacaagcg 329
Score = 22.3 bits (11), Expect = 4.6 Identities = 11/11 (100%)
 Strand = Plus / Minus
Query: 2192 gaaaaataact 2202
           Sbjct: 438 gaaaaataact 428
>AAV16677 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF
           06-JUN-1995 Polynucleotide sequence of a colon-specific
           gene. [Homo sapiens.]
         Length = 605
Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
Strand = Plus / Minus
Query: 2120 agcttcagcaa 2130
           пинийни
Sbjct: 312 agcttcagcaa 302
>AAV16674
           US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF
           06-JUN-1995 Polynucleotide sequence of a colon-specific
           gene. [Homo sapiens.]
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Length = 709



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Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Minus
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            11111111111
Sbjct: 201 tctttgttaat 191
>AAV16671 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF
          06-JUN-1995 Polynucleotide sequence of a colon-specific
          gene. [Homo sapiens.]
          Length = 548
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Minus
Query: 78 tgtttttcaca 88
          Sbjct: 19 tgtttttcaca 9
 Score = 22.3 bits (11), Expect = 4.6
Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 1106 tgttatgcacc 1116
            111111111
Sbjct: 108 tgttatgcacc 118
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 512 gctggtggcct 522
           Sbjct: 159 gctggtggcct 169
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 2146 agttcttaaag 2156
           ПППППП
Sbjct: 317 agttcttaaag 327
 Score = 22.3 bits (11), Expect = 4.6
 Identities = 11/11 (100%)
 Strand = Plus / Plus
Query: 1577 ggagatggagg 1587
            Sbjct: 372 ggagatggagg 382
  Database: us5733748.nt
   Posted date: Jan 24, 2002 1:37 PM
  Number of letters in database: 9317
  Number of sequences in database: 15
   bda K H
1.37 0.711
Lambda
                     1.31
Gapped
Lambda
   1.37 0.711
Matrix: blastn matrix:1 -3
Gap Penalties: Existence: 5, Extension: 2
Number of Hits to DB: 19
Number of Sequences: 15
Number of extensions: 19
Number of successful extensions: 19
Number of sequences better than 10.0: 8
length of query: 2609
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length of database: 9317

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effective HSP length: 12
effective length of query: 2597
effective length of database: 9137
effective search space: 23728789
effective search space used: 23728789
T: 0
A: 40
X1: 6 (11.9 bits)
X2: 15 (29.7 bits)
S1: 12 (24.3 bits)
S2: 11 (22.3 bits)
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us5733748.nt US5733748-A PR 06-JUN-1995 PF 06-JUN-1995 PCR primer used to amplify cloned colon-specific genes. >AAV16681 #HUMA−) HUMA**©** (Homo saniens.) ATGCTTCCGGCTCGTATG US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 PCR primer used to amplify cloned colon-specific genes. >AAV16682 [Homo sapiens.] GGGTTTTCCCAGTCACGAC >AAV16670 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom o sapiens.] ${\tt ATTGGTGCTACCTGGCTCTCTGTCTCTGCAGCTCTACAGGTGAGGCCCAGCAGAGGGAG}$ TAGGGCTCGCCATGTTTCTGGTGAGCCAATTTGGCTGATCTTGGGTGTCTGAACAGCTAT ${\tt TGGGTCCACCCCAGTCCCTTTCAGCTGCTGCTTAATGCCCTGCTCTCTCCCTGGCCCACC}$ ${\tt TTATAGAGAGCCCAAAGAGCTCCTGTAAGAGGGAGAACTCTATCTGTGGTTTATAATCTT}$ ${\tt ACCATGAGATCGGTCCAAGAGTCCATTCCCCAGGTGGGAGCCAACTGTCAGGGAGGTCTT}$ ${\tt TCCCACCAAACATCTTTCAGTTGCTGGGAGGTGACCATAGGGCTCTGCTTTTAAAGATAT}$ GGCTGCTTCAAAGGCCAGAGTCACAGGAAGGACTTCTTCCAGGGAGATTAGTGGTGATGG ${\tt AGAGGAGAGTTAAAATGACCTCATGTCCTTCTTGTCCACGGTTTTGTTGAGTTTTCACTC}$ TTCTAATGCAAGGGTCTCACACTGTGAACCACTTAGGATGTGATCACTTTCAGGTGGCCA GGAATGTTGAATGTCTTTGGCTCAGTTCATCTAAAAAAAGATATCTATTTGAAAGTTCTCA ${\tt TCACCAAGAGCCAATATCTAGGCATTTCCTCGGTAGCACAAATTTTCTNATTGCTTAGAA}$ AATTGTCCTCCTGTTCTTTCTGTCTGNAGACTTAAGTGAGTTAGGTCTTTAAGGAAAGC AACGCTCCTCTGAAATGCTTGTCTTTTTTCTGTTGCCGAAATAGCTGGTCCTTTTTCGGG AGTTAGATGTATAGAGTGTTTGTATGTAAACATTTCTTGTAGGCATCACCATGAACANAG ATATATTTCTATTTANTTANTATATGTGCACTTCAAGAAGTCACTGTCAGAGAATAAA GAATTGTCTTAAATGTCATGATTGGAGATGTCCTTTGCATTGCTTGGAAGGGGTGTACCT ${\tt AGAGCCAAGGAAATTGGCTCTGGTTTGGAAAAATTTTGCTGTTATTATAGTAAACATACA}$ AAGGATGTC >AAV16671 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom o sapiens.l ATGAGTCCTGTGAAAAACAATGTGGGCAGAGGCCTAAACATCGCCCTGGTGAATGGAACC ACGGGAGCTGTGCTGGGACAGAAGGCATTTGACATGTACTCTGGAGATGTTATGCACCTA GTGAAATTCCTTAAAGAAATTCCGGGGGGTGCACTGGTGCTGGTGGCCTCCTACGACGAT ${\tt CCAGGGACCAAAATGAACGATGAAAGCAGGAAACTCTTCTCTGACTTGGGGAGTTCCTAC}$ GCAAAACAACTGGGCTTCCGGGACAGCTGGGTCTTCATAGGAGCCAAAGACCTCAGGGGT ${\tt CCAGAGCTGCTGGAGATGGAGGGCTGCATGCCCCCGAAGCCATTTTAGGGTGGCTGTGGC}$ TCTTCCTCAGCCAGGGGCCTGAAGAAGYTCCTGCCTGCATTAGGAGTCANAGCCCGGCAG GCTGNAGGAGGAGCAGGGGGTGCTGCGTGGAAGGTGCTGCAGGCCTTGCACGCTGTG TCGCGCCT US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom >AAV16672 o sapiens.] TGTCTACTCAAGGTATTTCACAACTTATGACACGAATGGTAGATACAGTGTAAAAGTGCG GGCTCTGGGAGGAGTTAACGCAGCCAGACGGAGAGTGATACCCCAGCAGAGTGGAGCACT GTACATACCTGGCTGGATTGAGAATGATGAAATACAATGGAATCCACCAAGACCTGAAAT TAATAAGGATGATGTTCAACACAAGCAAGTGTGTTTCAGCAGAACATCCTCGGGAGGCTC ATTTGTGGCTTCTGATGTCCCAAATGCTCCCATACCTGATCTCTTCCCACCTGGCCAAAT CACCGACCTGAAGGCGGAAATTCACGGGGGCAGTCTCATTAATCTGACTTGGACAGCTCC TGGGGATGATTATGACCATGGAACAGCTCACAAGTATATCATTCGAATAAGTACAAGTAT TCTTGATCTCAGAGACAAGTTCAATGAATCTCTTCAAGTGAATACTACTGCTCTCATCCC AAAGGAAGCCAACTCTGAGGAAGTCTTTTTGTTTAAACCAGAAAACATTACTTTTGAAAA TGGCACAGATCTTTTCATTGCTATTCAGGCTGTTGATAAGGTCGATCTGAAATCAGAAAT ATCCAACATTGCACGAGTATCTTTGTTTATTCCTCCACAGACTCCGCCAGAGACACCTAG TCCTGATGAAACGTCTGCTCCTTGTGCCTAATATTCATATCAACAGCACCATTCCTGGCA TTCACATTTTAAAAATTATGTGGAAGTGGGTAGGAGAACTGCAGTTGTCAATAGNCTAGG GGTGAATTTTTGTGCGGTGAATAAATAATSATTTCANCCTTTTTTTGRTTTATAAAAAAA CGGNTNCCCATTGGGNNTNTNGNGGGGGGGGNNTTTTAA >AAV16673 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom o sapiens.1 AGTCGCTCTCCTAGCCCTTCTCTGTGCCTCACCCTCTGGCAATGCCATTCAGGCCAGGTC TTCCTCCTATAGTGGAGAGTATGGAGGTGGTGGAAAGCGATTCTCTCATTCTGGCAA CCAGTTGGACGCCCCATCACCGCCCTCCGGGTCCGAGTCAACACATACTACATCGTAGG GGAGGAGATCTTTCTGCACCCTGGGGAATCAGTGATCCAGGTTTCTGGGAAGTACAAGTG GTACCTGAAGAAGCTGGTATTTGTGACAGACAAGGGCCGCTATCTGTCTTTTGGGAAAGA CAGTGGCACAAGTTTCAATGCCGTCCCCTTGCACCCCAACACCGTGCTCCGCTTCATCAG TGGCCGGTCTGGTTCTCTCATCGATGCCATTGGCCTGCACTGGGATGTTTACCCCACTAG CTGCAGCAGATGCTGAGCCTCCTCTCCTTGGCAGGGGCACTGTGATGAGGAGTAAGAACT CCTTATCACTAACCCCCATC US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom >AAV16674 o sapiens.] TAAACTTGCTGTTTTGTTCCTGTGTCTTTGGTTGGTATTTCAGTAAGTTTTTGGT ATTCTCAAATTTTATCTAAATGGATAAACTATTAACATAGAACATAAACCCCAATTCTCC ATTTCATTTTCTCTTAGGCATGAATCATACAAAACTCAATATAGAGCAATGTTTGTAAT GAATTGTTCTATTAACAAAGAGGAGGTTCTAAGATATAAAGCCTCAGAGAACAGGAAGAA AAGGCGGGTCCATAAGAAGATGAGGTCTAACCGGGAAGATGCTGCTGAGAAGGCAGAGAC AGATGTGGAAGAAATCTATCACCCAGTCATGTGCACTGAATGTTCCACTGAAGTGGCAGT TTACGACAAGGATGAAGTCTTTCATTTTTTCAATGTTTTAGCAAGCCATTCCTAAACAGC CCAACTGGCATTTAATTACCCAATACTGTATATAAGGCAAATATGGACAGTTACTTTCCT

GTCGGGCTTTTTATTAAAGCTCTGTAGAATTAAAATGTTCTGGAATTAT >AAV16675 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom o sapiens.]



us5733748.nt



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GGTCATGGCTGCCAGAGCGCTCTGCATGCTGGGGCTGGTCCTGGCCTTGCTGTCCTCCAG
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TGCCCGGCTGTGATTGCTGCCAGGCACTGTTCATCTCAGCTTTTCTGTCCCTTTGCTCCC
GGAAGCGCTTCTGCTGAAAGTTCATATCTGGAGCCTGATGTTTAACGTAGTCCCATGCTC
САСССВААЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛ
          US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom
>AAV16676
o sapiens.]
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\tt CTGGGGCCATAACTCATGGGAAAACAGAATTATCCCCTAGGACTCCTTTCTAAGCCCCTA
>AAV16677
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o sapiens.l
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{\tt GGAGTCAAATATTCTGCCGAGCTTCACGTGGCTCACTGGAATTCTGCAAAGTACTCCAGC}
{\tt CTTGCTGAAGCTGCTCAAAGGCTGATGGTTTTGGCAGTTATTGGTGTTTTGATGAAGGTT}
GGTGAGGCCAACCCAAAGCTGCAGAAAGTACTTGATGCCCTCCAAGCAATTAAAACCAAG
TTCTGGACCTACCCTGGCTCTCTGACTCATCCTCCTCTTTATGAGAGTGTAACTTGGATC
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TCAAT
>AAV16678
          US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom
o sapiens.}
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GGTTTTGGGCATTGACACTGAGAGGCCCATTCTGGCAATGGACAGCTGTGTCTTTGCTGG
GGAGTATGAAGACACTCTAGGGACCTGTGTTATATTTGAAGAAAATGTTGAACATGCTGA
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ATGGCTGCAAATAAGGATATGGTTTCTCCCTTTGACCCAACAGGTTTGTTAACTTTTCTA
{\tt CCATGAAATTGAGGACGAGGAAGTGGTAGCTTTCAGCCCCGTTAAATCTTTGGATTTGGG}
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o sapiens.]
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>AAV16680
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o sapiens.)
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>AAV16668 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom o sapiens.]

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GGGGCCCGTGGTCCCACCTCTCCTGGTGTCTATGAAGCCTCTCTCCCGCTGCCCAGGA
CTACAGAATACCATCCCTGGTACCGTGTAGTTGCCGAAGTCCAGATCTGCCATGGCAAA
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GTGTCCGTGAGTCGTACACCTGATGGCTCCTCGTAGTCCGCCAGAAGGCAGGGGTCCAG
GTGTGGCTTGGAGACCAATGGAAAGTTGGAAAATTTGACGGGGAACCAATGATCTGGGATGATTCCC
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GCTTGWTCCAGTTCATCCCACCAGGAACGAAGGATTTT

>AAV16669 US5733748-A PA (HUMA-) HUMAN PR 06-JUN-1995 PF 06-JUN-1995 Polynucleotide sequence of a colon-specific gene. [Hom o sapiens]



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>DEX0075_1
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AGGTGTGTGTTTTTCACATCTTAAAGCTTATTCATTCCCAGGAACTCTTGGGGT
AGGTGTGTGTTTTTCACATCTTAAAGCTTAAAGCCTTGGGCCAGACAAATGTTCCATTCCTGAAGGAC $\tt CTCTCCAGAATCCGGATTGCTGAATCTTCCCTGTTGCCTAGAAGGGCTCCAAACCACCTCTTGACAATGG$ GTTTGTGGATGCCTTCCTGAAATATGAGAAGGCCGACAAATACTACTACAACAAGAAAAATCCTTGGGTCA ${\tt ACATTGGCCTGTGCCCGAGCGTCTGCTTCTGCTTGAATTTTAACAGCACGCTGATCCTGCTTCCTGTGT}$ GTCGCAATCTGCTGTCCTTCCTGAGGGGCACCTGCTCATTTTGCAGCCGCACACTGAGAAAGCAATTGGA CACCTGTTTAACTTTGACTGCTATAGCAGAAGCCGACAGGCCACAGATGGCTCCCTTGCCTCCATTCTCT CCAGCCTATCTCATGATGAGAAAAAGGGGGGTTCTTGGCTAAATCCCATCCAGTCCCGAAACACGACAGT GGAGTATGTGACATTCACCAGCGTTGCTGGTCTCACTGGAGTGATCATGACAATAGCCTTGATTCTCATG GTAACTTCAGCTACTGAGTTCATCCGGAGGAGTTATTTTGAAGTCTTCTGGTATACTCACCACCTTTTTA ${\tt TCTTCTATATCCTTGGCTTAGGGATTCACGGCATTGGTGGAATTGTCCGGGGTCAAACAGAGGAGGAGCAT}$ GAATGAGAGTCATCCTCGCAAGTGTGCAGAGTCTTTTGAGATGTGGGATGATCGTGACTCCCACTGTAGG $\tt CGCCCTAAGTTTGAAGGGCATCCCCCTGAGTCTTGGAAGTGGATCCTTGCACCGGTCATTCTTTATATCT$ GTGAAAGGATCCTCCGGTTTTACCGCTCCCAGCAGAAGGTTGTGATTACCAAGGTTGTTATGCACCCATC ${\tt CAAAGTTTTGGAATTGCAGATGAACAAGCGTGGCTTCAGCATGGAAGTGGGGCAGTATATCTTTGTTAAT}$ TGCCCCTCAATCTCTCCTGGAATGGCATCCTTTTACTTTGACCTCTGCTCCAGAGGAAGATTTCTTCT CCATTCATATCCGAGCAGCAGGGGACTGGACAGAAAATCTCATAAGGGCTTTCGAACAACAATATTCACC AATTCCCAGGATTGAAGTGGATGGTCCCTTTGGCACAGCCAGTGAGGATGTTTTCCAGTATGAAGTGGCT GTGCTGGTTGGAGCAGGAATTGGGGTCACCCCCTTTGCTTCTATCTTGAAATCCATCTGGTACAAATTCC AGTGTGCAGACCACAACCTCAAAACAAAAAAAGATCTATTTCTACTGGATCTGCAGGGAGACAGGTGCCTT TTCCTGGTTCAACAACCTGTTGACTTCCCTGGAACAGGAGGAGTGGGGAATTAGGCAAAGTGGGTTTTCTA AACTACCGTCTCTCCTCACCGGATGGGACAGCAATATTGTTGGTCATGCAGCATTAAACTTTGACAAGG CCACTGACATCGTGACAGGTCTGAAACAGAAAACCTCCTTTTGGGAGACCAATGTGGGACAATGAGTTTTTC TACAATAGCTACCTCCCACCCAAGTCTGTAGTGGGAGTTTTCTTATGTGGCCCTCGGACTTTGGCAAAG AGCCTGCGCAAATGCTGTCACCGATATTCCAGTCTGGATCCTAGAAAGGTTCAATTCTACTTCAACAAAG AAAATTTTTGAGTTATAGGAATAAGGACGGTAATCTGCATTTTGTCTCTTTTGTATCTTCAGTAATTGAGT TATAGGAATAAGGACGGTAATCTGCATTTTGTCTCTTTGTATCTTCAGTAATTTACTTGGTCTCNTCAGG TTTGANCAGTCACTTTAGGATAAGAATGTGCCTCTCAAGCCTTGACTCCCTGGTATTCTTTTTTTGATTG CATTCAACTTCGTTACTTGAGCTTCAGCAACTTAAGAACTTCTGAAGTTCTTAAAGTTCTGAANTTCTTA ${\tt AAGCCCATGGATCCTTTCTCAGAAAAATAACTGTAAATCTTTCTGGACAGCCATGACTGTAGCAAGGCTT}$ GATAGCAGAGTTTGGTGGTTCANAATTATACAACTAATCCCAGGTGATTTTATCAATTCCAGTGTTACC ATCTCCTGAGTTTTGGTTAATCTTTTGTCCCTCCCACCCCACAGAAGATTTTAAGTAGGGTGACTT CGAGAACCCCATCCCCATATAACACCAACAGTGTACATGTTTACTGTCACTTTTGATATGGTTTATCCAG ΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑ

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